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Community structure of woody plants on islands along a bioclimatic gradient

Paulo A.V. Borges^{1*}, Pedro Cardoso^{1,2*}, Simone Fattorini^{3*}, François Rigal^{1,4}, Thomas J. Matthews^{1,5,6}, Letizia Di Biase⁷, Isabel R. Amorim¹, Margarita Florencio^{1,8,9}, Luis Borda-de-Agua¹⁰, Carla Rego¹, Fernando Pereira¹, Rui Nunes¹, Rui Carvalho¹, Maria Teresa Ferreira¹, Heriberto López¹¹, Antonio J. Pérez Delgado^{11,12}, Rüdiger Otto¹³, Silvia Fernández Lugo¹³, Lea de Nascimento¹³, Juli Caujapé-Castells¹⁴, Juliane Casquet¹⁵, D P X H O ' D Q A Jacques Fournel¹⁷, Anne-Marie Sadeyen¹⁷, Rui B. Elias¹, José María Fernández-Palacios¹³, Pedro Oromí¹², Christophe Thébaud¹⁵, Dominique Strasberg¹⁷, Brent C. Emerson^{11,18}

*These authors contributed equally to this paper

- 1 and Environmental Changes Azorean Biodiversity Group and Universidade dos Açores, Rua Capitão João d'Ávila, São Pedro 9700-042 Angra do Heroísmo Terceira, Açores, Portugal.
- 2 Finnish Museum of Natural History University of Helsinki, P.O.Box 17, FIN-00014 Helsinki, Finland
- 3 Department of Life, Health & Environmental Sciences, University of L'Aquila, 67100 L'Aquila, Italy
- 4 CNRS-Université de Pau et des Pays de l'Adour, UMR 1213 Ecologie et Environnement, 64000 Pau, France
- 5 GEES (School of Geography, Earth & Atmospheric Sciences, The University of Birmingham, Birmingham B15 2TT, UK
- 6 Research, University of Birmingham, Edgbaston, Birmingham, B15 2TT, UK
- 7 Via Dante Alighieri 60, 00071, Pomezia, Italy
- 8 Departamento de Ecologia, Universidade Federal de Goiás, Avenida Esperança, s/n, Câmpus Samambaia, CEP 74690-900, Goiânia, Goiás, Brazil.
- 9 Department of Life Sciences, University of Alcalá, 28805 Alcalá de Henares, Madrid, Spain
- 10 REFER Biodiversity Chair, Universidade Nova de Lisboa, Av. da Universidade, 1700-055 Lisboa, Portugal.
- 11 Island Ecology and Evolution Research Group, IPNA-CSIC, C/ Carretera de la Laguna, 38206 La Laguna, Tenerife, Canary Islands, Spain.
- 12 Dept. of Animal Biology and Edaphology and Geology, University of Granada, 18014 Granada, Spain.
- 13 Instituto de Investigación en Recursos Cinegéticos, IREC (CSIC-UCLM-JCCM), Ronda de Toledo s/n, 13005 Ciudad Real, Spain.
- 14 Departament d'Ecologia i Evolució, Universitat de les Illes Balears, 07190 Esporles, Mallorca, Spain.
- 15 UMR 1166 Ecologie Evolutive et Fonctionnelle, CNRS, 91190 Brunoy, France.
- 16 UMR 1166 Ecologie Evolutive et Fonctionnelle, CNRS, 91190 Brunoy, France.
- 17 UMR 1166 Ecologie Evolutive et Fonctionnelle, CNRS, 91190 Brunoy, France.
- 18 UMR 1166 Ecologie Evolutive et Fonctionnelle, CNRS, 91190 Brunoy, France.

Abstract. Understanding patterns of community structure and the causes for their variation can be furthered by comparative biogeographic analyses of island biotas. We used woody plant data at the local scale to investigate variations in species rarity, alpha, beta, and gamma diversity within and between three islands from the oceanic archipelagoes of Azores, Canaries and Mascarene. We used standardized protocols to sample ten 50 m x 50 m forest plots in each of the three islands with contrasting climate and regional species pools: Terceira (Azores), Tenerife (Canaries), and Reunion (Mascarene Islands). Occupancy frequency distributions and species abundance distributions were used to investigate rarity. The partitioning of beta diversity in a distance-decay framework was used to test for spatial patterns of community composition. Rarity was much more pronounced in the highly diverse islands of Tenerife and Reunion than in the regionally poorer island of Terceira. The number of species rose faster with increasing sample area in both Tenerife and Reunion. The slope of the species rank abundance curve was steeper in Terceira whereas the richer island assemblages approached a lognormal model. Compositional changes according to spatial distance were mostly due to replacement of species in Terceira and Reunion. Our results point to important differences in the community structure of Terceira, which is the less diverse and temperate region in comparison to Tenerife and Reunion which are highly diverse.

Key words: Beta diversity partition, distance-decay, islands, rarity, species abundance distribution (SAD), species area relationship (SAR)

- ## Introduction

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Ten 50 m × 50 m square plots were established

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were implemented in R version 3.1.0 (R Core Team
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minimum and maximum number of species per plot
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restricted to indigenous species (i.e., endemics plot
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overall abundance.

and compared the following SAD models: the geometric series, the log-series, the Poisson lognormal

The lognormal is one of the most commonly used

Two main methods were employed to plot and model the species' abundances in which data were binned on rank one corresponds to the species with the highest abundance, rank two corresponding to the species with the second highest abundance, and so on. As a method to determine the best SAD model for any given

bins were constructed using base 100. The lowest AIC value was considered the best. Our nested SARs are similar to the

of beta diversity in Reunion, which can impact on the other two islands. Beta diversity in Reunion was the value of the slope, led us to restrict the analyses for Reunion to only four plots located within a 12 km than Tenerife. The density of woody plants in Terceira range, similar to the range covered by the 10 plots in Terceira and Tenerife. Thus, when we compared the slopes, between the three islands, we avoided

between plots in Reunion. The ten plots in Reunion. Alpha is the mean number of species in the ten plots. Max and Min alpha is the maximum and minimum number of species found in the ten plots, respectively. N is the number of individuals.

	Terceira	Tenerife	Reunion
Gamma	2186	715	1497
Alpha	7.3	7.4	25.6
Beta	0.51	1.16	1.61
Max alpha	9	9	33
Min alpha	5	5	16

the observed average distance was below the 0.025 and abundance data.

of wood plant rarity. In Terceira, only 18% of woody plant species occur in one plot whilst 45% occur in all ten plots (Tokeshi test). 38% of the species only occur in one plot whereas only 9% occur in all ten plots (Tokeshi test). 38% of the species only occur in one plot whereas only 9% occur in all ten plots (Tokeshi test).

regressed against the geographical distance between plots. The relationship between geographical distance and distance was tested using the R Package simba

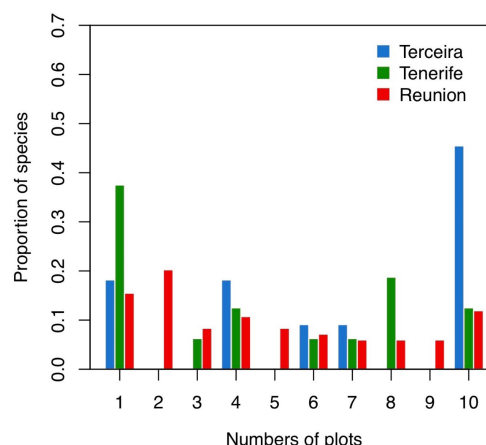


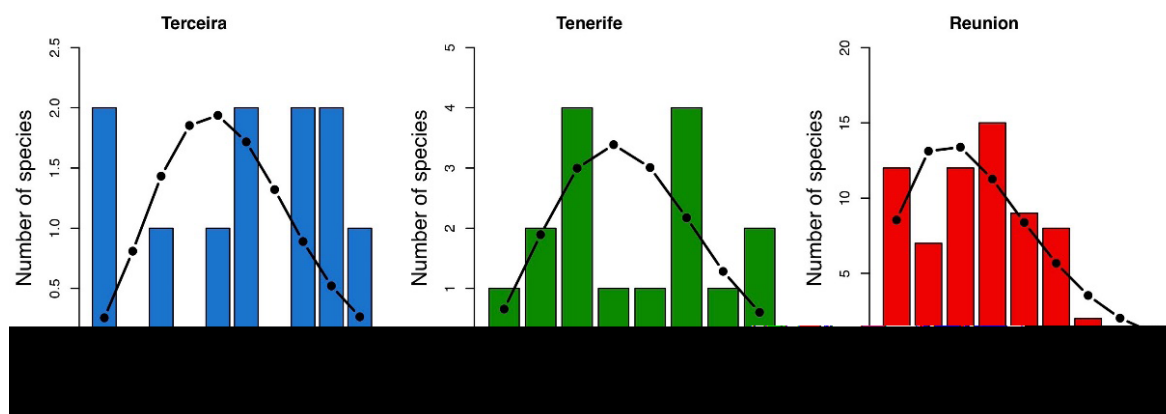
Figure 1. Proportion of species occurring in 1 to 10 plots for Terceira, Tenerife, and Reunion.

Results

Alpha, beta and gamma diversity patterns

93 woody plant species were sampled across thirty plots. Tenerife had very similar values of average alpha diversity

	Terceira	Tenerife	Reunion
Logseries	67.69	88.59	317.64
PLN	64.54	76	277.66
Gambin	65.46	75.31	274.36
Alpha	13.41	6.67	2.24



The following binning system was used: 0 corresponds to the number of species with 1 individual per species, bin 1 corresponds to the number of species with 2-3 individuals per species, bin 2 corresponds to the number of species

Species Area Relationships

The analyses of nested SARs confirmed the rapidity as sampling area increases in the two islands with the highest species richness (high α and β diversity) compared to the species poor and temperate island (low α and β diversity). The analyses of nested SARs confirmed the rapidity as sampling area increases in the two islands with the highest species richness (high α and β diversity) compared to the species poor and temperate island (low α and β diversity).

Figure 3. Geometric series rank-abundance plots for woody

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